SEQUENCE LISTING

HA	ASEGAWA, Mamoru IRONAKA, Takashi	
	aramyxoviral Vectors Encoding odies and Uses Thereof	
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	002-161964 002-06-03	
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tac act ctg agc agc tca gtg act gtc ccc tcc agc acc tgg ccc agc Tyr Thr Leu Ser Ser Ser Val Thr Val Pro Ser Ser Thr Trp Pro Ser 205 210 215	674
gag acc gtc acc tgc aac gtt gcc cac ccg gct tct agc acc aaa gtt Glu Thr Val Thr Cys Asn Val Ala His Pro Ala Ser Ser Thr Lys Val 220 225 230 235	722
gac aag aaa atc gta ccg cgc gac tgc taaccgtagt aagaaaaact Asp Lys Lys Ile Val Pro Arg Asp Cys 240	769
tagggtgaaa gttcatcgcg gccgtacggc c atg aaa caa agc act att gca Met Lys Gln Ser Thr Ile Ala 245 250	821
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tgg tat cag cgg aaa cag gga aaa tct cct cag ctc ctg atc tat ggt Trp Tyr Gln Arg Lys Gln Gly Lys Ser Pro Gln Leu Leu Ile Tyr Gly 300 305 310 315	1013
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Trp Thr Asp Gln Asp Ser Lys Asp Ser Thr Tyr Ser Met Ser Ser Thr
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ctc acg ttg acc aag gac gag tat gaa cga cat aac agc tat acc tgt
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gag gcc act cac aag aca tca act tca ccc att gtc aag agc ttc aac
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Glu Ala Thr His Lys Thr Ser Thr Ser Pro Ile Val Lys Ser Phe Asn
                                         470
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Val Arg Pro Gly Thr Ser Val Lys Ile Ser Cys Lys Ala Ser Gly Tyr
                            40
Thr Phe Thr Asn Tyr Trp Leu Gly Trp Val Lys Gln Arg Pro Gly His
Gly Leu Glu Trp Ile Gly Asp Ile Tyr Pro Gly Gly Gly Tyr Thr Asn
                                         75
                    70
Tyr Asn Glu Lys Phe Lys Gly Lys Ala Thr Leu Thr Ala Asp Thr Ser
                                    90
Ser Ser Thr Ala Tyr Met Gln Leu Ser Ser Leu Thr Ser Glu Asp Ser
            100
                                105
                                                     110
Ala Val Tyr Phe Cys Ala Arg Phe Tyr Tyr Gly Ser Ser Tyr Trp Tyr
                            120
        115
Phe Asp Val Trp Gly Gln Gly Thr Thr Val Thr Val Ser Ser Ala Lys
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                                             140
                        135
Thr Thr Pro Pro Ser Val Tyr Pro Leu Ala Pro Gly Ser Ala Ala Gln
                    150
                                        155
Thr Asn Ser Met Val Thr Leu Gly Cys Leu Val Lys Gly Tyr Phe Pro
                165
                                    170
Glu Pro Val Thr Val Thr Trp Asn Ser Gly Ser Leu Ser Ser Gly Val
            180
                                185
His Thr Phe Pro Ala Val Leu Gln Ser Asp Leu Tyr Thr Leu Ser Ser
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Ser Val Thr Val Pro Ser Ser Thr Trp Pro Ser Glu Thr Val Thr Cys
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215

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Pro Val Thr Lys Ala Asp Ile Glu Leu Thr Gln Ser Pro Ala Ile Met
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Ala Ala Ser Val Gly Glu Thr Val Thr Ile Thr Cys Gly Ala Ser Glu
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Asn Ile Tyr Gly Ala Leu Asn Trp Tyr Gln Arg Lys Gln Gly Lys Ser
Pro Gln Leu Leu Ile Tyr Gly Ala Thr Asn Leu Ala Asp Gly Met Ser
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Ser Arg Phe Ser Gly Ser Gly Ser Gly Arg Gln Tyr Ser Leu Lys Ile
                                    90
Ser Ser Leu His Pro Asp Asp Val Ala Thr Tyr Tyr Cys Gln Asn Val
                                105
                                                    110
Leu Ser Thr Pro Arg Thr Phe Gly Ala Gly Thr Lys Leu Glu Leu Lys
                            120
Arg Ala Asp Ala Ala Pro Thr Val Ser Ile Phe Pro Pro Ser Ser Glu
                                            140
Gln Leu Thr Ser Gly Gly Ala Ser Val Val Cys Phe Leu Asn Asn Phe
                    150
                                        155
Tyr Pro Lys Asp Ile Asn Val Lys Trp Lys Ile Asp Gly Ser Glu Arg
                                    170
Gln Asn Gly Val Leu Asn Ser Trp Thr Asp Gln Asp Ser Lys Asp Ser
            180
                                185
Thr Tyr Ser Met Ser Ser Thr Leu Thr Leu Thr Lys Asp Glu Tyr Glu
                            200
Arg His Asn Ser Tyr Thr Cys Glu Ala Thr His Lys Thr Ser Thr Ser
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Pro Ile Val Lys Ser Phe Asn Arg Asn Glu Cys
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230

225

Pro Arg Asp Cys

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ggctggtt
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<210> 16
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cagcgagacc
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caagagcttc aacaggaatg
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70
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<223> an anti-CD28 ScFv antibody gene (SYN205-13)
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qccaccatct cctqcaqaqc caqtqaqaqt gttqaatatt atqtcacaag tttaatgcag 120
tggtaccagc agaagccagg acagccaccc aaactcctca tctttgctgc atccaacgta 180
gaatetgggg teeetgeeag gtttagtgge agtgggtetg ggacaaactt cageeteaac 240
atccatcctg tggacgagga tgatgttgca atgtatttct gtcagcaaag taggaaggtt 300
ccttacacgt tcggaggggg gaccaagctg gaaataaaac ggggaggcgg cggttctggc 360
ggtggcggat caggtggcgg aggctcgcag gtgaaactgc agcagtctgg acctggcctg 420
gtgacgccct cacagagcct gtccatcact tgtactgtct ctgggttttc attaagcgac 480
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tgggctggtg gaggcacgaa ttataattcg gctctcatgt ccagaaagag catcagcaaa 600
gacaactcca agagccaagt tttcttaaaa atgaacagtc tgcaagctga tgacacagcc 660
gtgtattact gtgccagaga taagggatac teetattact attetatgga etaetgggge 720
caagggacca cggtcactgt ctcctcgtct aga
<210> 44
<211> 247
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Gln Arg Ala Thr Ile Ser Cys Arg Ala Ser Glu Ser Val Glu Tyr Tyr
Val Thr Ser Leu Met Gln Trp Tyr Gln Gln Lys Pro Gly Gln Pro Pro
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Lys Leu Leu Ile Phe Ala Ala Ser Asn Val Glu Ser Gly Val Pro Ala
Arg Phe Ser Gly Ser Gly Ser Gly Thr Asn Phe Ser Leu Asn Ile His
Pro Val Asp Glu Asp Asp Val Ala Met Tyr Phe Cys Gln Gln Ser Arg
Lys Val Pro Tyr Thr Phe Gly Gly Gly Thr Lys Leu Glu Ile Lys Arg
            100
                                105
Gly Gly Gly Ger Gly Gly Gly Gly Ser Gly Gly Gly Ser Gln
        115
                            120
                                                125
Val Lys Leu Gln Gln Ser Gly Pro Gly Leu Val Thr Pro Ser Gln Ser
   130
                        135
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Leu Ser Ile Thr Cys Thr Val Ser Gly Phe Ser Leu Ser Asp Tyr Gly
                    150
                                        155
Val His Trp Val Arg Gln Ser Pro Gly Gln Gly Leu Glu Trp Leu Gly
                165
                                    170
                                                         175
Val Ile Trp Ala Gly Gly Gly Thr Asn Tyr Asn Ser Ala Leu Met Ser
                                                    190
            180
                                185
Arg Lys Ser Ile Ser Lys Asp Asn Ser Lys Ser Gln Val Phe Leu Lys
                            200
        195
                                                205
Met Asn Ser Leu Gln Ala Asp Asp Thr Ala Val Tyr Tyr Cys Ala Arg
                        215
                                            220
Asp Lys Gly Tyr Ser Tyr Tyr Tyr Ser Met Asp Tyr Trp Gly Gln Gly
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                                        235
Thr Thr Val Thr Val Ser Ser
                245
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<211> 131
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<223> a NotI fragment containing an EIS sequence in
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qcqqccqcca aaqttcaatg gattttcagg tgcagatttt cagcttcctg ctaatcagtg 60
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atcgcggccg c
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Val Ile Met Ser Arg Gly
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gccaccatct
<210> 48
<211> 70
<212> DNA
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<220>
<223> a synthetic oligonucleotide for constructing an
      anti-CD28cst gene fragment
<400> 48
agggcagaga gccaccatct cctgcagagc cagtgagagt gttgaatatt atgtcacaag 60
tttaatgcag
<210> 49
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<212> DNA
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<223> a synthetic oligonucleotide for constructing an
      anti-CD28cst gene fragment
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tctttgctgc
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<211> 70
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                                                                    70
ggtggcggat
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acctggcctg
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gtgtattact
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